

GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:11:25 ; Search time 31 Seconds
(without alignments)
28686.456 Million cell
updates/sec

Title: US-09-300-482-225
Perfect score: 316
Sequence: 1
gataagggtgcgcacactgag.....gaatggagggagtacttcga 316

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2888648 seqs, 1391227531 residues

Total number of hits satisfying chosen parameters: 5777296

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_New:
1:
/ABSS/Data/CRF/ptodata/2/pna/PCT_NEW_COMB.seq:
2:
/ABSS/Data/CRF/ptodata/2/pna/US08_NEW_COMB.seq:
3:
/ABSS/Data/CRF/ptodata/2/pna/US09_NEW_COMB.seq:
4:
/ABSS/Data/CRF/ptodata/2/pna/US10_NEW_COMB.seq:
5:
/ABSS/Data/CRF/ptodata/2/pna/US11_NEW_COMB.seq:
6:
/ABSS/Data/CRF/ptodata/2/pna/US12_NEW_COMB.seq:
7:
/ABSS/Data/CRF/ptodata/2/pna/US12_NEW_COMB.seq1: